

**AMENDMENTS TO THE CLAIMS:**

Claims 26-29 are amended. The following is the status of the claims of the above-captioned application, as amended.

Claims 1-24 (Cancelled.)

Claim 25. (Currently amended.) A non-naturally occurring variant of a parent alpha-amylase, said variant comprising a substitution of a K at an amino acid position selected from the group consisting of 118, 320 and 458 (using SEQ ID NO:12 for numbering), wherein the variant has alpha-amylase activity, and wherein the variant has at least ~~80~~90% homology to SEQ ID NO:12.

Claim 26. (Currently amended.) The variant of claim 25, comprising an alteration ~~corresponding to~~of R118K.

Claim 27. (Currently amended.) The variant of claim 25, comprising an alteration ~~corresponding to~~of R320K.

Claim 28. (Currently amended.) The variant of claim 25, comprising an alteration ~~corresponding to~~of R458K.

Claim 29. (Currently amended.) The variant of claim 25, comprising an alteration ~~corresponding to~~of R118K, R320K and R458K.

Claim 30. (Previously presented.) The variant of claim 25, wherein the parent alpha-amylase is *Bacillus* sp. DSMZ no. 12649 alpha-amylase.

Claims 31-49 (Cancelled.)

Claim 50. (Previously presented.) The variant of claim 25, wherein the variant has at least 95% homology with an amino acid sequence of SEQ ID NO:12.

Claim 51-56. (Cancelled.)

Claim 57. (Previously presented.) The variant of claim 25, wherein the variant has at least 97% homology with an amino acid sequence of SEQ ID NO:12.

Claim 58. (Cancelled.)